



NCERT



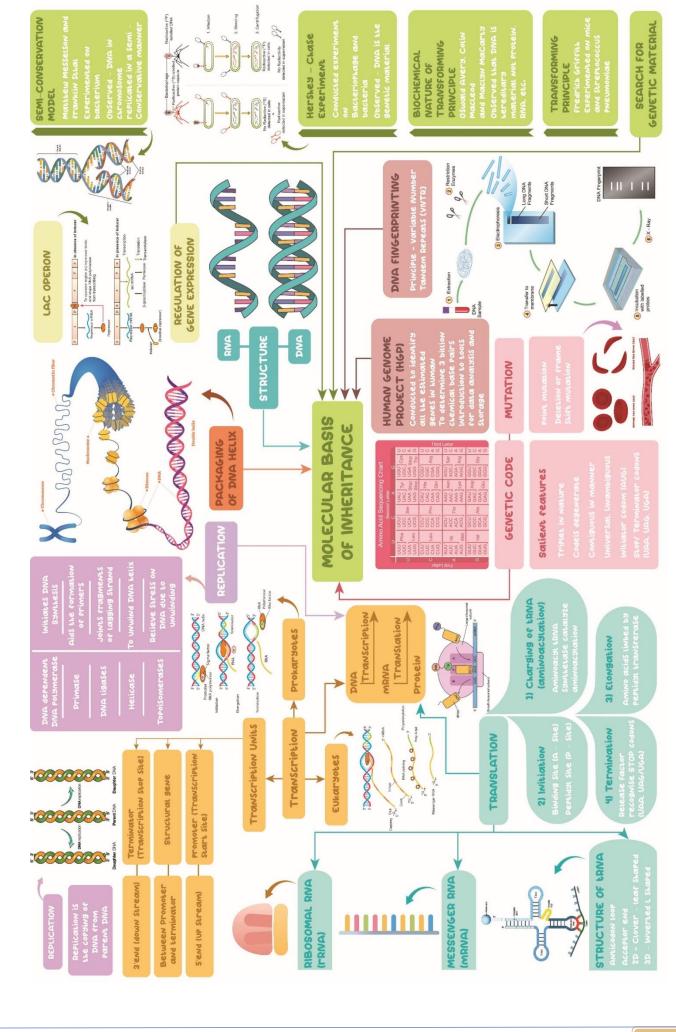
CHAPTER WISE TOPIC WISE

LINE BY LINE QUESTIONS





BY SCHOOL OF EDUCATORS



NCERT LINE BY LINE QUESTIONS

1.				acids are -	. 1	C) D:1	1 .1	D) 11 (11	
^		-	N T A	•	sides	C) Riboni	ıcleosides	D) None of these	
2.				types of -	مداده	C) Nucloi	a acida	D) Nucleamides	
	A) Nu	icieon	ues	B) Nucleo		C) Nuclei		D) Nucleamides	
					Pi	aragraph 6	.1		
2	Ŧ .:	1 (D	NTA .	11 1 6	1	The DNA			
3.	_			sually defin					
	•			eotides prese					
			_	of nucleotide	_	nt in it			
				pairs presen	it iii it				
4.	D) All			DNIA with	the corr	act arganism	C		
4.	Match	A	rigui oi	B B	me com	ect organism	.5 –		
	I) Φ ×			_	1.6 × 10e	bp (base pa	ire)		
	,		hage γ	,	3.3× 109	1 \	113)		
	III) E.	_	mage y		48502 bp	-			
			DNA (ł	naploid) 4) 5	_				
	I	II	III	IV					
	A) 4	3	1	2					
	B) 3	4	2	1					
	C) 4	3	1 2 2	1					
	D) 3	4	1	2					
					Pa	ragraph 6.1	1.1		
				Struct	ture of	polynucle	otides chair	ı	
5.	A nuc	leotid	es conta	ins –					
	A) He	exose s	ugar + 1	nitrogenous	base + p	phosphate gr	roup		
						ohosphate gr			
						ulphate grou			
					base +	sulphate gro	up		
6.			-	purine -					
	A) Cy								
	B) Ad								
	C) Thiamine D) More than one is correct								
7.	,			pyrimidine -	_				
, .				nymine		Guanine	D) None o	of these	
8.	•			•	,		,	or these	
		Which of these is a correct combination for a DNA nucleotides A) Oxyribose + Phosphate + Uracil							
	•	B) Oxyribose + Phosphate + Thymine							
		-		hosphate + i	•				
	D) De	oxyrib	ose + P	hosphate + '	Thymin	e			
9.		_		otides exists,	_				
	A) De					Both A & B	D) None o	of these	
10.	-	_	s base is			bon of pento	_		
44	A) 1′C		1 .	B) 2'C	C) 3		D) 5′C		
11.	Nitros	zenous	s base is	linked to p	entose s	ugar by which	ch bond –		

A) N – Glycosidic bond B) Phosphoester bond C) Phosphodiester bond D) Peptide bond Phosphate group is linked to which carbon of pentose sugar 12. B) 2'C D) 5′C 13. Identify the free ends of given polynucleotides chain -II A) 3' phosphate 5' hydroxyl B) 5' hydroxyl 3' phosphate C) 5' phosphate 3' hydroxyl D) 3' hydroxyl 5' phosphate 14. Backbone of polynucleotide chain is formed due to -A) Sugar and N-base B) Sugar and phosphate C) Phosphate and N - Base D) All of these 15. Which is correct about thymine & uracil -A) Uracil is 5-methyl thymine B) Thymine is 5-methyl uracil C) Uracil is 5-ethyl thymine D) Thymine in 5-ethyl uracil 16. DNA is -A) Acidic and positively charged B) Basic and positively charged C) Acidic and negatively charged D) Basic and negatively charged 17. Name of DNA as 'Nuclein' was given by -A) Francis crick B) Erwin Chargaff C) Friedrich Meischer D) Rosalind Franklin 18. Double Helix for structure of DNA model was proposed by -A) Wilkins and Franklin based on their Xray diffraction date B) Watson and Crick based on their X-ray diffraction date C) Chargaff based on their X-ray diffraction data D) None of these 19. The proposition of base pairing between the two stands of polynucleotide chain in double Helix model of DNA was based on observation of -A) Maurice Wilkins B) Rosalind Franklin D) Both A & B C) Erwin Chargaff

B) Anti-parallel polarity

D) Depends on organism

B) Peptide bond

D) Sulfide bond

20.

21.

22.

The two chains of double Helix DNA have -

The bases in two stands of DNA are paired through

Which of the following is true about base pairing in DNA -

A) Parallel polarity

A) Hydrogen bond

C) Glycosidic bond

C) No polarity

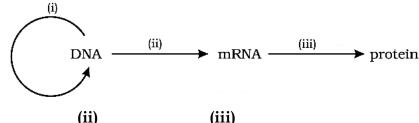
- A) Adenine forms two hydrogen bond with Guanine
- B) Adenine forms three hydrogen bond with Guanine
- C) Adenine forms two hydrogen bond with Thymine
- D) Adenine forms three hydrogen bond with Thymine
- 23. Which of the following is true about base pairing in DNA -
 - A) Guanine forms two H-bond with Cytosine
 - B) Guanine forms three H-bond with Cytosine
 - C) Guanine forms two H-bond with Adenine
 - D) Guanine forms three H-bond with Adenine
- Uniform distance between two stands of Helix is due to -24.
 - A) Double and triple bond formed between base pairs
 - B) Sugar phosphate backbone
 - C) Purine pyrimidine base pairing
 - D) None of these
- 25. How many of the following statements about Double - helix structure of DNA is correct
 - i) Two chains are coiled in right handed fashion
 - ii) Pitch of helix is 3.6 nm
 - iii) There are roughly 10 bp in each turn
 - iv) Plane of one base pair stacks over the other
 - A) 1
- B) 2
- C)3

D) 4

- 26. Pitch of helix in double helix DNA is -
 - A) 3.6 nm
- B) 3.4 nm
- C) 3.2 nm

- D) 3.8 nm
- Central dogma in molecular biology was proposed by --27.
 - A) Crick
- B) Watson
- C) F. Meischar
- D) Chragaff

28. Identify correct labels --



- (i)
- Translation
- (iii)

- A) Replication B) Replication
- Transcription
- Transcription Translation

- C) Transcription
- Replication
- Translation

- D) Translation
- Replication
- Transcription

Paragraph 6.1.2 **Packaging of DNA Helix**

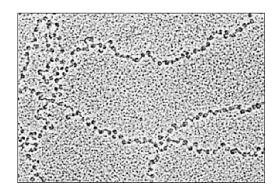
- 29. If length of E. coli DNA is 1.36 mm, calculate number of base pair in E. coli? Given - distance between consecutive base pairs is 0.34×10^{-9} m. -
 - A) 4×106
- B) 4×109
- C) $4 \times 10-6$
- D) 4×1012
- **Assertion**: In E. coli, DNA is scattered throughout the cell 30.

Reason: In E. coli, there is no defined nucleus

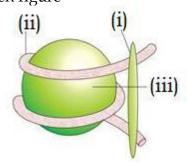
- A) Both Assertion & Reason are correct & Reason is correct explanation for Assertion
- B) Both Assertion & Reason are correct but Reason is not correct explanation for Assertion
- C) Assertion is correct and Reason is incorrect
- D) Reason is correct and Assertion is incorrect
- 31. Histones are -
 - A) Positive and acidic in eukaryotes
 - B) Positive and acidic in prokaryotes

- C) Positive and basic in eukaryotes
- D) Positive and basic in prokaryotes
- Assertion Histones are positively charged Reason Histones are rich in basic amino acid 32. residues lysine and arginine
 - A) Assertion & Reason are correct and Reason is correct explanation for Assertion
 - B) Assertion & Reason are correct and Reason is not the correct explanation for Assertion
 - C) Assertion is correct and Reason is wrong
 - D) Both Assertion and Reason are wrong
- 33. Histones are organized into-
 - A) Hexamer
 - B) Octamer C) Tetramer
- D) Dimer
- 34. A typical nucleosome contain _____ bp of DNA has
 - A) 200
- B) 400
- C) 600
- D) 800

- 35. Repeating unit of chromatin -
 - A) Are nucleosomes
 - B) Are seen as 'beads-on-string' under electron microscope
 - C) Are packed to form fibres
 - D) All of these
- 36. The figure show -



- A) Beads-on-string
- B) A nucleosome
- C) Chromatin
- D) More than one option is correct
- 37. Identify the correct label for given figure



- (i)
- (ii)

(iii)

- A) H2 histone **DNA**
 - Histone octamer DNA
- Histone octamer

- B) H1 histone C) H2 histone
- Histon octamer
- DNA

- D) H1 histone
- **DNA**
- Histone octamer
- 38. Chromosomes are connected chromatin fibres present -
 - A) At all times in cell
 - B) Only during cell division formed at prophase
 - C) Only during cell division formed at metaphase
 - D) Only during cell division formed at Interphase

39.	<i>J</i> 1		omatin are present. C	Choose the correct set of		
	characters for heterochromat					
	i) Loosely packed					
	iii) Light stain i	•	•			
	v) Inactive chromatin			D)		
4.0	· · · · · · · · · · · · · · · · · · ·	, vi	C) i, iii, vi	D) ii, iv, v		
40.						
	i) Loosely packed					
	iii) Light stain					
	v) Inactive chromatin					
	A) i, iii, v B) ii, iv, vi		D) ii, iv, v			
			aterial Transform	ning Principle		
41.	Griffith's experiments were c					
	A) 1928 B) 1958	C) 1978	D) 1968			
42.	The experiment of Griffith w	_				
	A) Diplococcus pneumoniae,		B) Haemophilus ir	ıfluenzas, fungi		
	C) Streptococcus pneumonia	e, fungi	D) None of these			
43.	O					
	I II		II			
	` '	` '	lucous coat			
	ii. S-strain (b) Rough col		o mucous coat			
	A) (i)-(a)-(1) B) (i)-(b)-(1)			2)		
44.	Which strain of the microbe t					
	A) S-strain B) R-str	,	,			
45.	Griffith observed that the mid	e died surprising	ly the following com	bination of strains was used		
	which was unusual-		5) 1.11 1.0			
	A) S-strain heat killed	0	B) Heat killed S-str			
4.6	C) Heat killed R-strain + Live	S-strain	D) Heat killed S-st	rain + Live R-strain		
46.	In Griffith experiment					
	A) R-strain transformed to S-					
	B) R-strain transformed to S-s					
	C) S-strain transformed to R-					
4.77	D) S-strain transformed to R-	strain and lost viri	ulence			
47.	Griffith claimed that-	1 1 .				
	A) Some protein was transfer	· ·	ıa			
	B) Some DNA was transferre	O	1			
	C) Some carbohydrates was transferred among bacteria					
	D) None of these		C.TT. C .	D : 1		
4.0			of Transforming	-		
48.	Prior to work of Avery, Macl	-	_	_		
40	A) Protein B) DNA	,	NA	D) None		
49.	A very, Macleod & McCarty					
	A) DNA caused transformati					
	B) RNA caused transformation					
	C) Protein caused transforma					
- 0	D) Lipid caused transformati					
50.	Which enzyme inhibited the		N T	D) A11		
	A) Protease B) RNa	se C) D	Nase	D) All		
	6.2.	l The Genetic N	Material is DNA			

51. Unequivocal proof that DNA is genetic material came from experiments of-A) Avery, Macleod & McCarty B) Hershey and Chase C) de Vries, Correns and Tschermak D) Sutton and Boveri 52. The scientists of Q-11 worked with-C) a fungi A) a virus B) a bacteria D) a nematode 53. In the experiment performed for proving DNA as genetic material, the bacteriophages were grown on medium containing-A) radioactive sulfur B) radioactive nitrogen C) radioactive phosphorous D) More than one option 54. The bacteriophages growing in presence of radioactive phosphorous __(i)__ contained radioactive __(ii)__. (i) (ii) A) P32 **DNA** B) P35 Protein C) P32 Protein D) P35 **DNA** 55. Bacteriophages grown on radioactive Sulphur __(i)__ contained radioactive__(ii)__. (i) (ii) **DNA** A) S32 B) S32 Protein C) S35 DNA D) P35 Protein 56. The bacteria involved in Hershey & chase experiment of 1952 was-A) Bacteriophage B) E. coli C) S. pneumoniae D) C. butyliwm Bacteria infected with virus that showed radioactivity had-57. B) radioactive DNA (S35) A) radioactive DNA (S32) C) radioactive DNA (P32) D) radioactive DNA (P35) 58. Bacteriophage Radioactive (³²P) Radioactive (³⁵S) labelled labelled DNA protein capsule 2. 3. No Radioactive (35S) Radioactive (³²P) detected in cells detected in cells Radioactive (³⁵S) No Radioactivity & detected in supernatant detected in supernatant

Identify the correct label.

A) Blending Infection Centrifugati on

B) Infection Blending Centrifugati on C) Centrifugati on Infection **Blending** D) Blending Infection Centrifugati on 6.2.2 Properties of Genetic Material (DNA vs RNA) RNA is genetic material in-B) QB Bacteriophage A) TMV C) Both A and B D) None of these Properties of genetic material include-A) Stable B) Mutable C) Replicable D) All of these A - Stability as a property of genetic material was very evident in Griffith's transforming principle. R-Heat can kill the bacteria and completely destroy the properties of genetic material A) Both A and R are true and R is correct explanation for A B) Both A and R are true but R is not correct explanation for R C) A is true but R is false D) Both A and R are false Which is more structurally and chemically stable? A) DNA B) RNA D) All C) Protein RNA viruses show-A) Less mutation B) Faster evolution C) Slower evolution D) More than one option is correct DNA is preferred by nature over RNA for-A) Storage of genetic information B) Transmission of genetic information C) Expression of genetic information D) More than one 6.3 RNA World Choose incorrect statement RNA-A) was first genetic material B) acts as catalyst too C) is more stable than DNA D) has protein synthesizing mechanism built around it 6.4 Replication Scheme for replication of DNA was proposed by -B) Meselson & Stahl

A) Watson & Crick C) Taylor D) Hershey & Chase

The replication of DNA is-67.

> A) Conservative B) Non-conservative

C) Semi-conservative D) All of these depending on organism

68.

59.

60.

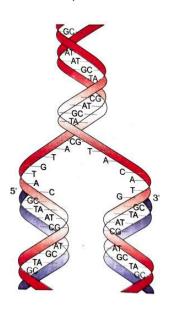
61.

62.

63.

64.

65.

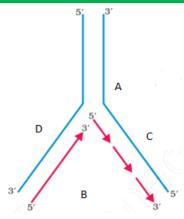


The figure shows-

A) Conservative DNA replication model B) Semi-conservative DNA replication model C) Non-conservative DNA replication model D) Can't say 6.4.1 The Experimental Proof 69. The DNA replication model experimental proof was first shown in-A) Human cells B) E. coli C) Plant cell D) Vicia faba 70. Meselson and stahl performed experiment for proving DNA replication scheme in-A) 1952 B) 1953 C) 1958 D) 1961 71. The bacteria were grown in medium containing-A) 15NH₄Cl - 15N is heavy isotope of nitrogen B) 14NH₄Cl – 14N is heavy isotope of nitrogen C) 15NH₄Cl – 15N is normal isotope of nitrogen D) 14NH₄Cl - 14N is normal isotope of nitrogen 72. The heavy DNA molecule containing heavy isotope of N is distinguished from normal DNA by-A) UV rays B) Ethidium bromide solution C) Centrifugation is CsCl density gradient D) PCR technique In Meselson & Stahl experiment, first they-73. A) grew bacteria on heavy isotope of N medium followed by normal one B) grew bacteria on normal isotope of N medium followed by heavy one C) grew bacteria on radioactive N followed by Keavy one D) grew bacteria on heavy isotope of N followed by radioactive one 74. Identify the correct label Generation I 15 N-DNA N-DNA -(i) 14 N 15 N Hybrid Light Heavy (i) (iii) (ii) A) Light Heavy Hybrid B) Heavy Hybrid Light C) Light Hybrid Light D) Heavy Heavy Hybrid 75. In Meselson & Stahl expt a bacteria after dividing in 20 minutes had a hybrid DNA. What will be the ratio of Hybrid to Light after 80 minutes? A) 2:14B) 14:2 C) 16:2D) 2:16 Similar experiment on Vicia faba was conducted by _____ to detect distribution of newly 76. synthesized DNA in chromosomes. A) Taylor B) Stahl C) Gamow D) Nirenberg 77. Experiment on Vicia faba involved use of-A) Radioactive uridine B) Radioactive thymidine C) Radioactive adenosine D) Radioactive cytidine 6.4.2 The Machinery and the Enzymes

78. The main enzyme of replication is-

	A) RNA dependent RNA polymerase
	B) RNA dependent DNA polymerase
	C) DNA dependent DNA polymerase
	D) DNA dependent RNA polymerase
79.	Choose correct statement with regard with efficiency of DNA polymerase.
	A) 4.6 × 106 bp of E. coli replicate within 46 minutes
	B) The average rate of polymerization of DNA polymerase has to be approximately 2000
	bp/minute
	C) The polymerization accuracy is very high and very fast
	D) All of these
80.	What is function of deoxyribonucleoside triphosphate -
	A) It act as substrate
	B) Provide energy for polymerization
	C) A and B both
	D) It is product formed after polymerization
81.	Assertion: The two strands of DNA cannot be separated in their length.
01.	Reason: Separation required very high energy.
	A) Both Assertion & Reason are correct and reason is correct explanation of assertion
	B) Both Assertion & Reason are correct and reason is not correct explanation of assertion
	C) Assertion is correct, Reason is false
	D) Assertion & Reason are false
82.	Polymerization by DNA polymerase is in-
-	A) $3' \rightarrow 5'$ direction only
	B) $5' \rightarrow 3'$ direction only
	C) A and B both
	D) Random
83.	The template of replication fork with polarity $5' \rightarrow 3'$ is while $3' \rightarrow 5'$ is
	A) continuous, continuous
	B) continuous, discontinuous
	C) discontinuous, continuous
	D) discontinuous, discontinuous
84.	DNA ligase act on-
	A) $5' \rightarrow 3'$ template strand
	B) $3' \rightarrow 5'$ template strand
	C) Both A and B
	D) Ligate RNA with vector of $3' \rightarrow 5'$ polarity
85.	The replication is eukaryotes take place in-
	A) M-phase B) G1 phase C) S-phase D) G2 phase
86.	Polyploidy resulted by-
	A) A failure in cell division after DNA replication
	B) A failure in DNA replication after cell division
	C) A failure in cell division before DNA replication
	D) A and C both
87.	Correct label of A, B, C, D is-



- (i) A = Template parental strand
- (ii) B = Newly synthesized strand
- (iii) D = Continuous strand
- (iv) C = Discontinuous strand
- A) i, ii only
- B) iii, iv only
- C) i, ii, iii, iv
- D) None of these

6.5 Transcription

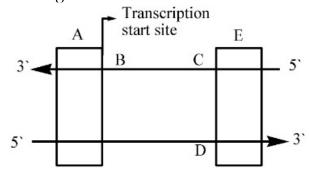
- 88. Transcription is-
 - A) The process of copying genetic information from both strand of DNA into RNA
 - B) The process of copying genetic information from one strand of DNA into RNA
 - C) The process of copying genetic information from RNA into DNA
 - D) A and B both
- 89. In transcription, adenosine bind with
 - A) Thymine
- B) Uracil
- C) Cytosine
- D) A and B both
- 90. Why both the strand of DNA are not copied during transcription-
 - A) If both strands act a template, they would code for RNA molecules with same sequence
 - B) RNA formed by transcription of both strand, when code for protein, the sequence of amino acid in protein are same
 - C) The two RNA molecules if produced simultaneously would be complementary to each other
 - D) All of these
- 91. Translation of RNA would be prevented if-
 - A) RNA is single strand
 - B) RNA is double-stranded
 - C) RNA is produced by both strand of DNA
 - D) B and C both

6.5.1 Transcription Unit

- 92. Transcription unit primarily consist of-
 - A) 1 region
- B) 2 regions
- C) 3 regions
- D) None of these

- 93. Transcription primarily required-
 - A) RNA dependent RNA polymerase
 - B) DNA dependent RNA polymerase
 - C) DNA dependent DNA polymerase
 - D) RNA dependents DNA polymerase
- 94. Template strand of transcription unit is/are-
 - A) $5' \rightarrow 3'$ strand of DNA
 - B) $3' \rightarrow 5'$ strand of DNA
 - C) Site of catalysis of enzyme required for transcription
 - D) B and C both
- 95. What is coding strand of given template strand 3' AGCATGCA 5'
 - A) 5' TACGTACGT 3'

- B) 5' UACGUACGU 3'
- C) 3' UACGUACGU 5'
- D) 3' TACGTACGT 5'
- 96. Label A, B, C, D, E of given diagram.



	a	b	С	d	e
A	Promoter	Structural	Template	Coding	Terminator
		Gene	strand	strand	
В	Terminator	Structural	Coding	Template	Promoter
		Gene	strand	strand	
C	Promoter	Template	Coding	Structural	Terminator
		strand	strand	Gene	
D	None of these				

- 97. Promoter is located-
 - (i) 3' end
 - (ii) 5' end
 - (iii) upstream of structural gene
 - (iv) downstream of structural gene
 - A) i, iii
- B) ii, iii

- C) i, iv
- D) ii, iv

- 98. Terminator is located at
 - (i) 3' end
 - (ii) 5' end
 - (iii) upstream of structural gene
 - (iv) downstream of structural gene
 - A) i, iii
- B) ii, iii

- C) i, iv
- D) ii, iv

6.5.2 Transcription unit and the gene

- 99. A gene is defined as-
 - A) Functional unit of inheritance
 - B) Non-functional region of DNA that haven't any information
 - C) A and B both
 - D) None of these
- 100. Cistron is-
 - A) Segment of DNA coding for a polypeptide
 - B) Segment of RNA coding for a polypeptide
 - C) Segment of DNA that are non-coding sequence
 - D) Segment of RNA have not any coding sequence
- 101. Choose the correct statement.
 - A) Monocistronic eukaryotic structural gene have interrupted coding sequence.
 - B) Polycistronic prokaryotic structural gene have interrupted coding sequence.

	C) Monocistronic prokaryotic structural gene have interrupted coding sequence.
	D) A and B both
102.	Exons are-
	A) Coding sequence
	B) Non-coding sequence
	C) Expressed sequence
	D) A and C both
102	,
103.	Intron-
	A) appear is mature or processed RNA
	B) do not appear in mature or processed RNA
	C) appear is prokaryotes
	D) B and C both
	6.5.3 Types of RNA & Process of Transcription
104.	Which of following play role is protein synthesis of prokaryote?
	A) r-RNA B) t-RNA
	C) m-RNA D) All of these
105.	The function of some RNA are given below choose the incorrect one.
	A) mRNA provide template strand
	B) mRNA provide non-template strand
	C) tRNA bring amino acid
	D) rRNA play structural & catalytic role
106	How many polymerase required is bacteria for transcription of all type of RNA?
100.	
107	, , , , , , , , , , , , , , , , , , , ,
107.	Choose incorrect step about transcription.
	A) RNA polymerase binds to promoter and initiate transcription.
	B) Nucleotide triphosphate act as substrate and polymerization in a template.
	C) A short stretch of RNA remains bound to enzyme.
400	D) Last step is termination.
108.	Initiation factor and termination factor are-
	A) Sigma and Rho factor respectively.
	B) Rho and Sigma factor respectively.
	C) Rho and Rho factor respectively.
	D) Sigma and Sigma factor respectively.
109.	Translation & transcription in eukaryote occur in
	A) Cytoplasm & nucleus respectively
	B) Nucleus & cytoplasm respectively
	C) Cytosol
	D) Nucleus
110.	Which of following can be coupled in bacteria?
	A) Replication & transcription
	B) Transcription & translation
	C) Replication & translation
	D) None of these
111.	Transcription of 18 s rRNA is done byin eukaryote.
	A) RNA pol. I B) RNA pol. II C) RNA pol. III D) All of these
112.	Choose incorrect statement.
	A) 5.8 s r-RNA and 5 s-RNA transcribes by same RNA polymerase in eukaryote.
	B) hnRNA & mRNA transcribe by same RNA polymerase in eukaryote.
	C) tRNA & snRNA transcribes by same RNA polymerase in eukaryote.
	D) None of these
	Dy Notic of diese

113.	Splicing is rec	•					
	•	tron in eukaryote					
	•	on in eukaryote					
	,	on in prokaryote					
444	•	tron in prokaryote					
114.	Capping is-	((1 1)	1				
	•	• •	triphosphate at 5' end				
	B) addition of adenylate residue at 3' end C) addition of methyl guanosine triphosphate at 3' end						
	•	•					
115	•	f adenylate residue a	t 5 end				
115.		cessed hnRNA is- B) mRNA	C) rRNA	D) None of these			
	A) tixiA	b) ilixivA	6.6 Genetic Code	•			
116.	George Gamo	ow argued-	0.0 Genetic Code				
	~	<u> </u>	ev have code for 20 a	mino acid the code should constitute a			
	combination of	=	J				
	B) There are o	only 4 bases and if the	ey have code for 20 ar	mino acid the code should constitute a			
	combination of	2					
	C) Genetic co	de is triplet					
	D) B and C	-					
117.	Which of foll	owing have maximu	m codon in genetic co	ode-			
	A) Leu	B) Met	C) Cal	D) Phe			
118.		owing is/are showing	_				
	A) UUU	B) AUG	C) UGA	D) GUA			
119.		of codon is for-	C) C1	D) 61			
	A) Met	B) Phe	C) Gls	D) Gly			
12 0	0.11 11		utations and Gene	etic Code			
120.		nemia is classical exai	mple ot-				
	A) point muta						
	B) frameshift						
	C) deletion m						
101	D) addition m	nnaemia, there are ch	anges in gone for				
141.	A) alpha glob		anges in gene for-				
	B) beta globir						
	C) gamma glo						
	D) delta globi						
122.			fect of mutation is cha	ange of amino acid residue-			
	A) Valine to a			O			
	B) Valine to g						
	C) Alanine to						
	D) Glutamic a	acid to valine					
123.	_	_	G RED CAP ⇒ BIG RE	EM DCA P			
	A) Deletion n						
	B) Point muta						
	C) Addition r						
104	D) More than	_	CAD				
124.		$AP \Rightarrow RAM HAS BIC$	ı CAP				
	The given exa	_					
	A) Addition r	nutation					

- B) Deletion mutation
- C) Substitution mutation
- D) More than one option

6.6.2 tRNA - the Adapter Molecule

- 125. tRNA has-
 - A) Codon loop
- B) Anticodon loop
- C) Both
- D) Neither
- 126. The presence of adapter molecule to read the code on DNA and bind to amino acids was postulated by-
 - A) James Watson
- B) Francis Crick
- C) Friedrich Meisher
- D) Both A and B

- 127. tRNA was also called-
 - A) s RNA (soluble RNA)

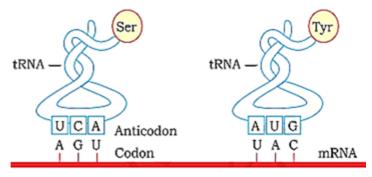
B) s RNA (single RNA)

C) s RNA (smart RNA)

- D) s RNA (simple RNA)
- 128. The amino acid acceptor end of tRNA is-
 - A) 5'

- B) 3'
- C) Can be any of these
- D) Free end

- 129. For initiation translation,
 - A) Only tRNA carries initiator amino acid to the site.
 - B) Specific rRNA carries initiator amino acid to the site.
 - C) Any rRNA carries initiator amino acid to the site.
 - D) Specific tRNA carries initiator amino acid to the site.
- 130. For stop codon-
 - A) There are specific tRNAs with amino acids.
 - B) There are specific tRNAs which do not bind to any amino acids.
 - C) There are no tRNA.
 - D) There are tRNAs which may or may not bind to amino acids.
- 131. The given figure shows-



- A) Secondary structure of tRNA Cloverleaf
- B) Primary structure of tRNA clover-leaf
- C) Secondary structure of tRNA -inverted-L
- D) Primary structure of tRNA inverted-L

6.7 TRANSLATION

- 132. Translation refers to process of-
 - A) Making RNA from DNA
 - B) Making DNA from RNA
 - C) Polymerization of nucleotide to form a DNA
 - D) Polymerization of amino acid to form a polypeptide
- 133. The order and sequence of amino acid during translation are defined by-
 - A) The sequences of bases in r-RNA
 - B) The sequences of bases in t-RNA
 - C) The sequences of bases in m-RNA
 - D) All of these
- 134. Which of following bond is formed during translation?

- A) Glycosidic bond
- B) Phosphodiester bond
- C) Peptide bond
- D) All of these
- 135. First phase of translation does not involve -
 - A) Charging of RNA
 - B) Amino acids are activated in presence of ATP
 - C) Activated amino acid are linked to their cognate tRNA
 - D) None of these
- 136. Initiation or first phase of translation is -
 - A) Amino acylation of tRNA
 - B) Amino acylation of mRNA
 - C) Both A and B
 - D) Deamino acylation of mRNA
- 137. The cellular factory responsible for synthesizing protein is-
 - A) Ribosome B) Lysosome
- C) Peroxisome
- D) None of these
- 138. In inactive state, protein factory of cell exist in
 - A) Two state

- B) 4 state in prokaryote
- C) 6 state in eukaryote
- D) B and C both
- 139. Which of following is sign as beginning of translation?
 - A) When the large subunit of protein factory of cell encounters an mRNA.
 - B) When the small subunit of protein factory of cell encounters an mRNA.
 - C) When the small subunit of protein factory of cell encounter a tRNA.
 - D) When the large subunit of protein factory of cell encounters a tRNA.
- 140. The bond formation (peptide) between charged tRNA is accomplished due to-
 - A) Presence of ATP and catalyst
 - B) Two such charged tRNA are brought close by two site in large subunit of ribosome
 - C) Two charged tRNA are brought close by two site in small subunit of ribosome & presence of ATP along with catalyst
 - D) A and B both
- 141. The ribosome act as catalyst during bond formation (peptide) as in-
 - A) 28 s rRNA in bacteria

- B) 23 s rRNA in bacteria
- C) 23 s rRNA in eukaryote
- D) 28 s tRNA in bacteria
- 142. Choose the correct statement-
 - A) A translational unit in mRNA is sequence of RNA that is flanked by start codon and stop codon and codes for polypeptide.
 - B) A translational unit is sequence of DNA that is flanked by start codon & codes for polypeptide.
 - C) A transcriptional unit in tRNA is the sequence of RNA that is flanked by start codon and stop codon and codes for polypeptide.
 - D) A transcriptional unit in rRNA is the sequence of RNA that is flanked by start codon (AUG) and stop codon and codes for polypeptide.
- 143. UTR is/are -
 - (i) Untranslated region of mRNA
 - (ii) It present at both 5' end (start codon) and 3' end (before stop codon)
 - (iii) They are required for efficient translation process
 - (iv) It present at both 3' end (before start codon) and 3' end (after stop codon)
 - A) i, ii, iii are correct
- B) i, ii, iii and iv are correct
- C) i, iii, iv are correct
- D) i, iii are correct
- 144. Initiator tRNA binds with

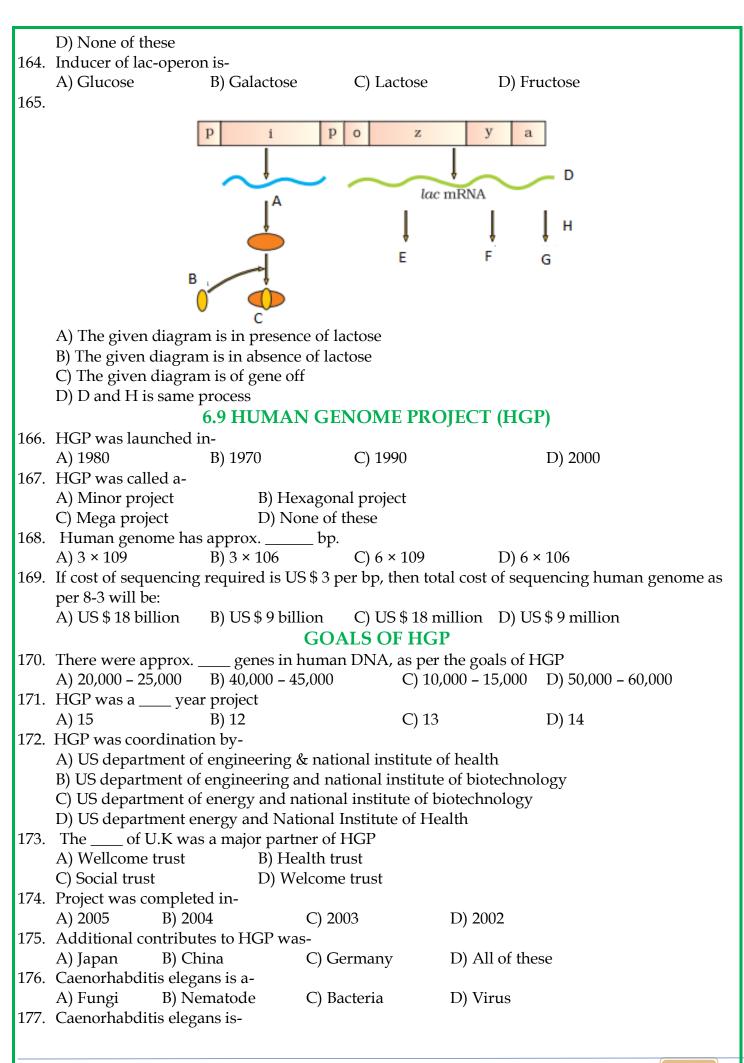
- A) AUG codon of mRNA
- B) at initiation of protein synthesis
- C) ATG codon of dsDNA
- D) A and B both
- 145. Choose the correct about elongation of translation-
 - A) Complexes composed of an amino acid linked to tRNA, sequentially bind to appropriate codon in mRNA by forming complementary base pairs with the tRNA anticodon
 - B) The ribosome moves from codon to codon along the mRNA in $(3' \rightarrow 5')$.
 - C) Complexes composed of an amino acid linked to tRNA, sequentially bind to appropriate anticodon in mRNA by forming complementary base pair with tRNA codon.
 - D) A and B both
- 146. Termination of translation complex is done when-
 - A) Release factor binds with stop codon (AUG)
 - B) Release factor binds with UGA like codon
 - C) Complete translation of DNA including UTR occurs in eukaryotes
 - D) B and C both

6.8 REGULATION OF GENE EXPRESSION

BOTANY

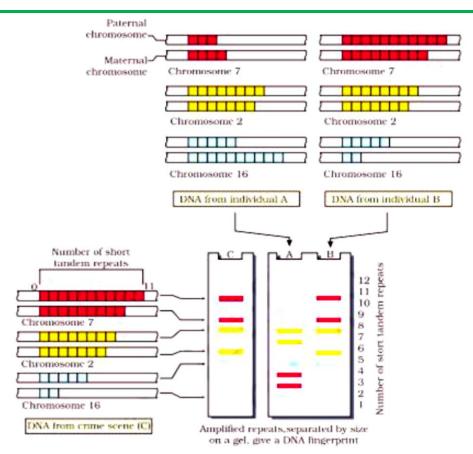
- 147. Gene regulation is eukaryote exerted at-
 - A) Formation of primary transcript
 - B) Transport of mRNA from nucleus to cytoplasm
 - C) A and B both
 - D) Regulation of splicing of tRNA
- 148. β -glactosidase is used to catalyze the hydrosis of
 - A) Lactose into galactose and glucose
 - B) Lactose into fructose & glucose
 - C) Lactose into fructose & fructose
 - D) None of these
- 149. E.coli do not have lactose around them to utilized for energy source, they would-
 - A) No longer require the synthesis of enzyme α -galactosidase
 - B) Synthesized enzyme β -galactosidase
 - C) Die due to lack of carbon source and energy source
 - D) None of these
- 150. In prokaryote, predominant site for control of gene expression is-
 - A) Control of rate of transcriptional initiation
 - B) Control of rate of translational
 - C) Control of rate of transcriptional elongation
 - D) B and C both
- 151. Given below are statement. Choose the incorrect statement.
 - A) The development and differentiation of embryo into adult organisms are result of coordinated regulation of expression of several sets of genes.
 - B) Regulatory proteins act positively in activator.
 - C) In a transcriptional unit the activity of RNA polymerase at a given promoter is in turn regulated by interaction with accessory protein.
 - D) None of these
- 152. Operator-
 - A) Region adjacent to sequence by which repressor mRNA formed
 - B) Bind with repressor protein
 - C) Bind with inducer
 - D) A and B both

153.	Each operon h	as -					
	A) Same operator and same repressor						
	B) Same operator but specific repressor						
	C) Specific ope	erator b	ut same r	epressor			
	D) Specific ope			_			
				6.8.1 THE LAC OI	PERON		
154.	Lac operon wa		ed first by	y-			
	A) François Jac						
	B) Jacque Mon		1				
	C) Geneticist a		chemist				
155	D) None of the						
155.	Lac operon is/		1 1			0	
						er & regulators genes.	
				e is regulated by a con			
		_	патогу де	ene, monocistronic su	ucturar gene nav	ving five gene along with	
	promotor & op D) A and C bo						
156	Regulatory gen		-operon i	C_			
150.	A) p-gene	B) i-ge		c) o-gene	D) z-gene		
157	i in i-gene star		ric .	C) 0-gene	D) Z-gene		
157.	A) inducer		hitor	C) A and B both	D) Inactive r	enressor	
158	Match Column	,		C) II and D both	D) mactive i	cpressor	
100.	Column-I(Ger			olumn-II(Product)			
	(a) Z-gene) Repressor mRNA			
	(b) a-gene) β-gal			
	(c) Y-gene			i) Permease			
	(d) i-gene) Transacetylase			
	a b	c	d `	,			
	A) (iii) (ii)	(iv)	(i)				
	B) (iv) (iii)		(i)				
		(iv)	(iii)				
	D) ii) (iv)	(iii)	(i)				
159.	The monomer:	ic produ	uct of lact	ose is chiefly hydroly	zed by-		
	A) i-gene		B) z-gene	e C) a-gene	D) y-8	gene.	
160.	Lac in lac-oper		r				
	A) Monosacch			B) Disaccharide			
	C) Polysacchar			D) Insect			
161.		_		_	_	th medium of bacteria, th	
		-		through by action of p	•	5	
1.0	A) i-gene		B) z-gene	e C) a-gene	D) y-8	gene	
162.	Allolactose is-	1					
	A) Inducer of I	_					
	B) Inductive re	_		the man decate of monages	ou ma DNIA amadii	alailait tuomaaninatian af	
	•		it bina wi	th product of repress	or mikina and ii	nhibit transcription of	
	structural gene						
162	D) All of these						
103.	Lac operon is A) Negative re		n oneron				
	B) Positive reg	_	_				
	C) A and B bo		operon				
	-, and - 00						



A) Free living , non-pathogenic B) Parasitic, pathogenic C) Free living , pathogenic D) Parasitic, non-pathogenic 178. Methods / approaches of HGP include-A) Excess sequence tags B) Expressed sequence tags C) Exercise sequence tags D) Exerted sequence tags 179. Sequence annotations refer to-A) Identifying all genes expressed as RNA and then sequencing then B) Sequencing the whole set of genome and then assigning different regions with functions C) Identifying and sequencing the genome simultaneously D) More than one option is correct 180. For sequencing, the DNA is-A) Partially extracted from cell B) Totally isolated from cell C) Not needed to isolated from cell D) None of these 181. The DNA for sequencing is converted to fragments of small size. The fragments are made-A) On a pre - decided basis B) On a pre - defined basis C) Randomly D) Depending upon organism 182. The step in DNA sequencing after fragmentation of DNA is-A) Cloning in host using vectors B) Cloning in vectors using host C) Amplification of DNA fragments D) More than one option 183. Commonly used hosts for DNA cloning include-B) BAC A) Bacteria C) YAC D) Both A and C 184. BAC stands for-A) Bacterial artificial colour B) Binominal artificial characterization C) Bacterial artificial chromosome D) Bacterial articular chromosome 185. Fragments were sequenced using automated DNA sequence that worked on principle of a method developed by-A) Erwin Chargaff B) Marshal Nirenberg C) Frederick Sanger D) George Gamow 186. Method for determination of amines acid sequence in protein was developed by-A) Erwin Chargaff B) Marshal Nirenberg C) Frederick Sanger D) George Gamow 187. The last of the 24 human chromosomes to be sequenced was-B) Chromosome X A) Chromosome 1 D) Chromosome Y C) Chromosome 22 6.9.1 SALIENT FEATURES OF HUMAN GENOME 188. According to HGP, human genome contains-A) ~ 3000 million bp B) ~ 6000 million bp C) \sim 9000 million bp D) ~ 1000 million bp

189.	Dystrophin was found to be-		
	A) Largest known human gene with	n 2.4 million bases	
	B) Smallest known human gene with		
	C) Largest known human gene with		
	D) Smallest known human gene wit		
190.	Which chromosome was found to ha		
	A) Chr 22 B) Chr 1		D) Chr Y
191.	Which chromosome was found to h	ave fewest genes-	•
	A) Chr X B) Chr Y		D) Chr 5
	•	IA FINGER PRINTING	G
192.	The DNA sequence in which small s		
		B) Repetitive DNA	3
	,	D) More than one option	
193.	Satellite DNA classified into differen	· -	atellite, mini-satellite, etc based
	on-	O	,
	A) Length of segment	B) Number of repetitive	
	, e	D) All of these	
194.	Polymorphism arises due to-	,	
	A) Mutation – inheritable	B) Stability of geneti	ic material
	C) Mutation - non-heritable	D) All of these	
195.	DNA polymorphism is observed mo	ore in-	
	A) non-coding DNA sequence as its	mutation affects reproduc	etion
	B) coding DNA sequence as its muta	ation affects reproduction	
	C) non-coding DNA sequence as it i	mutation may not affect rej	production ability
	D) Coding DNA sequence as its mu	tation may not affect repro	duction ability
196.	Technique of DNA fingerprinting w	as initially developed by-	
	A) James Watson B) Jansl	ley	
	C) Alec Jeffreys D) Mah	neshwari	
197.	VNTR stands for-		
	A) Various number of Tendon Repe		
	B) Variable Number of Tendon Repo		
	C) Various Number of Tandem Rep	eats	
	D) Variable Number of Tandem Rep	peats	
198.	VNTR belongs to-		
	•	B) Macro-satellite	
	C) Mini-satellite	D) All of these	
199.			



In the given figure if 'C' is the DNA collected from crime site and 'A' & 'B' are samples from suspects, than who is the criminal?

A) B

B) A

C) Both A and B

D) None of these

NEET PREVIOUS YEARS QUESTIONS

- 1. The experimental proof for semi-conservative replication of DNA was first shown in a: [2018]
 - (a) Fungus
- (b) Bacterium
- (c) Virus
- (d) Plant

2. Select the correct match

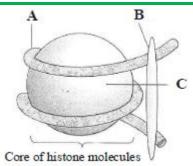
(a) Alec Jeffreys

- Streptococcus pneumoniae - TMV
- (b) Alfred Hershey and Martha Chase
- (c) François Jacob and Jacques Monod
- Lac operon
- (d) Matthew Meselson and F. Stahl
- Pisum sativum
- 3. Many ribosomes may associate with a single mRNA to form multiple copies of a polypeptide simultaneously. Such strings of ribosomes are termed as: [2018]
- (b) Polyhedral bodies (c) Nucleosome
- (d) Plastidome
- 4. AGGTATCGCAT is a sequence from the coding strand of a gene. What will be the corresponding sequence of the transcribed mRNA? [2018]
 - (a) AGGUAUCGCAU (b) UGGTUTCGCAT (c) UCCAUAGCGUA

(d) ACCUAUGCGAU [2018]

[2018]

- 5. All of the following are part of an operon except:
- (d) An enhancer
- (a) An operator (b) Structural genes (c) A promoter The given figure shows the structure of nucleosome with their parts labelled as A, B & C. Identify A, B 6. and C. [2017]



(a) A – DNA; B – H1 histone; C – Histone octamer (b) A – H1 histone; B – DNA; C – Histone octamer (c) A – Histone octamer; B – RNA; C – H1 histone (d) A – RNA; B – H1 histone; C – Histone octamer Match the codons given in column I with their respective amino acids given in column II and choose the correct answer. [2017]

ľ		Column -I		Column -II
		(Codons)		(Amino acids)
	A	UUU	I.	Serine
	В	GGG	II.	Methionine
	C	UCU	Ш.	Phenylalanine
	D	CCC	IV.	Glycine
	E	AUG	V.	Proline

- (a) A III; B IV; C I; D V; E II(b) A - III; B - I; C - IV; D - V; E - II
- (c) A III; B IV; C V; D I; E II(d) A - II; B - IV; C - I; D - V; E - III
- 8. The final proof for DNA as the genetic material came from the experiments of: [2017]
- (a) Hershey and Chase

7.

- (b) Avery, Mcleod and McCarty (c) Hargobind Khorana (d) Griffith
- 9. DNA fragments are: [2017]
- (a) Negatively charged (b) Neutral
 - (c) Either positively or negatively charged depending on their size
 - (d) Positively charged
- Which of the following RNAs should be most abundant in animal cell? [2017] 10.
- (a) t-RNA (b) m-RNA (c) mi-RNA 11. The association of histone H₁ with a nucleosome indicates
 - [2017]
- (a) DNA replication is occurring.
 - (b) the DNA is condensed into a chromatin fibre.
- (c) the DNA double helix is exposed. (d) transcription is occurring.
- If there are 999 bases in an RNA that codes for a protein with 333 amino acids, and the base at position 12. 901 is deleted such that the length of the RNA becomes 998 bases, how many codons will be altered? [2017]
 - (a) 11 (b) 33 (c) 333 (d) 1
- During DNA replication, Okazaki fragments are used to elongate 13. [2017]
 - (a) the lagging strand towards replication fork. (b) the leading strand away from replication fork.
- (c) the lagging strand away from the replication fork. (d) the leading strand towards replication fork. [2017]
- Spliceosomes are not found in cells of: 14. (a) Fungi (b) Animals (d) Plants (c) Bacteria
- **15.** Which of the following is required as inducer(s) for the expression of Lac operon? [2016]
- (b) Galactose (c) Lactose (d) Lactose and galactose (a) Glucose
- Which of the following is not required for any of the techniques of DNA fingerprinting available at 16. present? [2016]
 - (a) Polymerase chain reaction
- (b) Zinc finger analysis

(c) Restriction enzymes

- (d) DNA-DNA hybridisation
- **17.** Which one of the following is the starter codon?

[2016]

- (a) AUG
- (b) UGA
- (c) UAA
- (d) UAG

(d) r-RNA

18.	A complex of ribosomes atta	ched to a single	e strand of RNA is	known as:		[2016]
	(a) Polysome (b) Po	_	(c) Polypeptide		(d) Okazaki fra	agment
19.	There are three genes a, b, c.	Percentage of	crossing over betw	een a and b	is 20%, b and c	is 28% and a
	and c is 8%. What is the sequ	uence of genes	on chromosome?			[2015]
	(a) b, a, c (b) a,		(c) a, c, b	(d) No	ne of these	
20.	Which one of the following is					[2015]
	(a) 5' phosphoryl and 3' hydr	oxyl ends	(b) Heterocyclic	_	bases	
	(c) Chargaff's rule		(d) Watson and C			
21.	In sea urchin DNA, which is				•	
	percentages of the other three	-			e:	[2015]
	(a) G-17%, A-16.5%, T-32.5		(b) G-17%, A-33	•	0.7	
22	(c) G-8.5%, A-50%, T-24.5%		(d) G-34%, A-24		%	[2017]
22.	The movement of a gene from				 •	[2015]
22	\ / 1	nslocation	(c) crossing over	\ /	rersion	[2015]
23.	Gene regulation governing la					: [2015]
	(a) negative and inducible be	*				
	(b) negative and repressible l(c) feedback inhibition becau					
	(d) positive and inducible be		_		ascription.	
24.	Satellite DNA is important b		madeed by factose	•		[2015]
27.	(a) shows high degree of pol		nonulation and also	the same d	egree of nolym	
	individual, which is heritable			o the same a	egree or porymo	orphism m un
	(b) does not code for protein	_		ne populatio	n.	
	(c) codes for enzymes needed			re populario		
	(d) codes for proteins needed					
25.	Select the correct option.	,				[2014]
	Direction of RNA	Direction	f reading of			. ,
	synthesis		e DNA strand			
	·		e DNA stranti			
	(a) 5'—3'	3′—5′				
	(b) 3'—5'	5'—3' 5'—3'				
	(c) 5'—3'	l				
	(d) 3'—5'	3′—5′				
2.0	(a) 5'—3' 3'—5' (b) 3'-				_5′3′ <u>_</u> 5′	1004.43
26.	Which one of the following is	represents a pa				[2014]
	(a) 5' - GAATTC - 3'		(b) 5' - CCAATG			
	3' - CTTAAG - 5'		3' - GATACC			
	(c) 5' - CATTAG - 3' 3' - GATAAC - 5'		(d) 5' - GATACC	- 3		
27.	Transformation was discover	red by ·				[2014]
<i>4</i> 1 •	(a) Meselson and Stahl (b) H		ase (c) Griffith	(d) W ₂	tson and Crick	[2017]
28.	Which one of the following	•	` /	(d) 110	itson and Crick	[2014]
-0.			n from DNA to tRN	VA.		[2011]
			in mRNA to make			
	(c) Repressor protein – Bind					
			perator and promot			
29.	Commonly used vectors for					[2014]
	(a) T-DNA (b) BAC and YA	_		(d) T/A	A cloning vector	
30.	Purines found both in DNA	` /		. /	-	[NEET-2019
	(1) Adenine and thymine					
	(3) Guanine and cytosine					
31.	Under which of the following	g conditions w	ill there be no chan	ge in the rea	ading frame of f	
	mRNA?	_				[NEET-2019
		5'AAC	AGCGGUGCUAU	JU3'		

	(1) Insertion of G at 5th position (2) Deletion of G from 5th position	
	(3) Insertion of A and G and 4th and 5th positions respectively	
	(4) Deletion of GGU from 7th, 8th and 9th positions	
32.	Which of the following features of genetic code does allow bacteria to produce hu	
	recombinant DNA technology?	[NEET-2019]
	 (1) Genetic code is not ambiguous (2) Genetic code is redundant (3) Genetic code is nearly universal (4) Genetic code is specific 	
		DVTTT 40401
33.	Expressed Sequence Tags (ESTs) refers to :-	[NEET-2019]
	(1) Genes expressed as RNA (2) Polypeptide expression	
2.4	(3) DNA polymorphism (4) Novel DNA sequences	DIEDE 20101
34.	Match the following genes of the Lac operon with their respective products:	[NEET-2019]
	(a) i gene (i) b-galactosidase (b) z gene (ii) Permease	
	(c) a gene (iii) Repressor (d) y gene (iv) Transacetylase Select the correct option.	
	(a) (b) (c) (d) (a) (b) (c) (d) (a) (b) (c) (d) (a) (b) (c) (d)	(d)
	(a) (b) (c) (d) (a) (b) (c) (d) (d) (d) (d) (d) (e) (d) (d) (e) (d) (d) (e) (d) (e) (d) (e) (d) (e) (d) (e) (e) (e) (e) (e) (e) (e) (e) (e) (e	
35.	What will be the sequence of mRNA produced by the following stretch of DNA?	(11)
55.		7-2019 ODISSA]
	3'ATGCATGCATGCTG' TEMPLATE STRAND	2017 ODISSN
	5' TACGTACGTACGTAC3' CODING STRAND	
	(1) 3'AUGCAUGCAUGCS' (2) 5'UACGUACGUACGUAC 3'	
	(3) 3' UACGUACGUACGUAC 5' (4) 5' AUGCAUGCAUGCAUG 3'	
36.		Γ-2019 ODISSA]
	(a) RNA polymerase I (i) tRNA	,
	(b) RNA polymerase II (ii) rRNA	
	(c) RNA polymerase III (iii) hnRNA	
	Select the correct option from the following:	
	(1) a-i, b-iii, c-ii (2) a-i, b-ii, c-iii (3) a-ii, b-iii, c-i	
37.	From the following, identify the correct combination of salient features of Genetic	
		[-2019 ODISSA]
	(1) Universal, Non-ambiguous, Overlapping (2) Degenerate, Overlapping, Comma	
20	(3) Universal, Ambiguous, Degenerate (4) Degenerate, Non-overlapping, No	
38.	Which scientist experimentally proved that DNA is the sole genetic material in bac	1 0
	•	[-2019 ODISSA]
20		ob and Monod
39.	In the process of transcription in Eukaryotes, the RNA polymerase I transcribes [N (1) mRNA with additional processing, capping and tailing (2) tRNA, 5 S rRNA and	
	(3) rRNAs.28 S, 18 S and 5.8 S (4) Precursor of mRN	
40.	What initiation and termination factors are involved in transcription in Eukaryotes	
+ ∪.		[-2019 ODISSA]
	(1) s and r, respectively (2) a and b, respectively	1-2017 ODISSA _j
	(3) b and g, respectively (4) a and s, respectively	
41.		T-2020 COVID]
	(1) Franklin (2) Meischer (3) Chargaff (4) Mendel	1
42.	In the polynucleotide chain of DNA, a nitrogenous base is linked to the -OH of:	NEET-2020 COVID
	(1) 2'C pentose sugar (2) 3'C pentose sugar (3) 5'C pentose sugar (4) 1'C pentose	
43.	E.coli has only 4.6×10^6 base pairs and completes the process of replication within	18 minutes; then
	the average rate of polymerisation is approximately-	[-2020 COVID]
	(1) 2000 base pairs/second (2) 3000 base pairs/second	
	(3) 4000 base pairs/second (4) 1000 base pairs/second	
44.	Which is the basis of genetic mapping of human genome as well as DNA finger pr	
		T-2020 COVID]
	(1) Polymorphism in DNA sequence (2) Single nucleotide polymorphism	
	(3) Polymorphism in hnRNA sequence (4) Polymorphism in RNA sequence	

45.	Name the enzyme that facilitates opening of DNA helix du			[NEET-2020]
4.6	1) RNA polymerase 2) DNA ligase 3) DNA helic	ase 4) DNA p	oolymerase	DIEDE AGAIL
46.	The first phase of translation is: 1) Recognition of an anti–codon 2) Binding of mRNA	to ribosomo		[NEET-2020]
	3) Recognition of DNA molecule 4) Aminoacylation of			
47.	If the distance between two consecutive base pairs is 0.34 DNA double helix in a typical mammalian cell is 6.6×10 ⁹	nm and the total m		
	approximately	op, men me lengm	of the DNA	[NEET-2020]
	1) 2.7 meters 2) 2.0 meters 3) 2.5 meters	4) 2.2 me	ters	
48.	Complete the flow chart on central dogma			[NEET-2021]
	(a) \bigcirc DNA $\xrightarrow{\text{(b)}}$ mRNA $\xrightarrow{\text{(c)}}$ (d)			
	1) (a)-Translation; (b)-Replication; (c)-Transcription;(d)-T	'ransduction		
	2) (a)- Replication; (b)- Transcription; (c)- Translation; (d)-F			
	3) (a)-Transduction; (b)-Transcription (c)-Transduction; (D			
	4) a) Replication, b) Transcription, c) Transduction; D)-pro			
49.	Identify the correct statement			[NEET-2021]
	1) RNA polymerase binds with Rho factor to terminate the		iption in bact	teria.
	2) The coding strand in a transcription unit is copied to an	mKNA		
	3) Split gene arrangement is characteristic of prokaryotes4) In capping, methyl guanosine triphosphate is added to tl	ne 3' and of hnDN	٨	
50.	What is the role of RNA polymerase III in the process of ti			[NEET2021]
50.	± *	nscribes precursor	-	[TELETZOZI]
		nscribes rRNAs (2		5.8S)
51.	Which is the "Only enzyme" that has "capability" to cataly	yse initiation, Elon	gation and To	ermination in
	the process of transcription in prokaryotes?			[NEET-2021]
	1. DNA dependent RNA polymerase 2. DNA Liga	se		
	1 1 0	endent DNA polym	erase	
52.	A specific recognition sequence identified by endonuclease			ions within the
	DNA is			[NEET-2021]
	· · · · · · · · · · · · · · · · · · ·	omic Nucleotide s	_	
50		erate primer seque	ice	DIEEE 20211
53.	Which of the following RNAs is not required for the synth 1) tRNA 2) rRNA 3) siRNA) mRNA	[NEET-2021]
54.	1) tRNA 2) rRNA 3) siRNA Statement-I: The codon 'AUG codes for methionine and ph		IIIKINA	[NEET-2021]
57.	Statement-II: AAA' and 'AAG" both codons code for the a			[14221]
	In the light of the above statements, choose the correct ans	•	ns given belo	ow
	1) Both Statement I and Statement II are false	_	_	
	2) Statement I is correct but Statement II is false			
	3) Statement I is incorrect but Statement II is true			
55	4) Both Statement I and Statement II are true	··· · · · · · · · · · · · · · · · · ·		INFET 20201
55.	Which one of the following statements about Histones is w 1) The pH of histones is slightly acidic.	rong:		[NEET-2020]
	2) Histones are rich in amino acids – Lysine and Arginine.			
	3) Histones carry positive charge in the side chain			
	4) Histone are organized to form a unit of 8 molecules.			
56.	DNA fingerprinting involves identifying differences in sor	ne specific regions	in DNA sequ	uence, called
	as:			[NEET-2021]
	, 1	ymorphic DNA	,	llite DNA
57.	In an <i>E.coli</i> strain <i>i</i> gene gets mutated and its product		nducer mole	
	growth medium is provided with lactose, what will be	the outcome?		[NEET-2022]
	1) Only z gene will get transcribed			

- 2) z, y, a genes will be transcribed
- 3) z, y, a genes will not be translated
- 4) RNA polymerase will bind the promoter region
- If the length of a DNA molecule is 1.1 metres, what will be the approximate number of base 58. pairs? [NEET-2022]
 - 1) 3.3×10^9 bp
- 2) 6.6×10^9 bp 3) 3.3×10^6 bp
- 4) 6.6×10^6 bp
- Ten *E.coli* cells with ¹⁵ N dsDNA are incubated in medium containing ¹⁴ N nucleotide. After 60 59. minutes, how many E.coli cells will have DNA totally free from ^{15}N ? [NEET-2022]
 - 1) 20 cells
- 2) 40 cells
- 3) 60 cells
- 4) 80 cells

NCERT LINE BY LINE QUESTIONS - ANSWERS

	-> -								
1) D	2) C	3) D	4) A	5) B	6) B	7) B	8) D	9) D	10) A
11) A	12) D	13) C	14) B	15) B	16) C	17) C	18) A	19) C	20) B
21) A	22) C	23) B	24) C	25) C	26) B	27) A	28) B	29) A	30) D
31) C	32) A	33) B	34) A	35) D	36) D	37) D	38) C	39) D	40) C
41) A	42) D	43) C	44) A	45) D	46) A	47) D	48) A	49) A	50) C
51) B	52) A	53) D	54) A	55) D	56) B	57) C	58) B	59) C	60) D
61) C	62) A	63) B	64) A	65) C	66) A	67) C	68) B	69) B	70) C
71) A	72) C	73) A	74) C	75) A	76) A	77) B	78) C	79) C	80) C
81) A	82) B	83) C	84) A	85) C	86) A	87) C	88) B	89) B	90) C
91) D	92) C	93) B	94) D	95) A	96) A	97) B	98) C	99) A	100) A
101) C	102) D	103) B	104) D	105) B	106) A	107) B	108) A	109) A	110) B
111) A	112) A	113) A	114) A	115) B	116) B	117) A	118) B	119) A	120) A
121) B	122) D	123) D	124) A	125) B	126) B	127) A	128) B	129) D	130) C
131) A	132) D	133) C	134) C	135) D	136) A	137) A	138) A	139) B	140) D
141) B	142) A	143) D	144) D	145) A	146) B	147) C	148) A	149) D	150) A
151) D	152) B	153) D	154) C	155) B	156) B	157) B	158) D	159) B	160) B
161) D	162) A	163) C	164) C	165) A	166) C	167) C	168) A	169) B	170) A
171) C	172) D	173) A	174) C	175) D	176) B	177) A	178) B	179) B	180) B
181) C	182) D	183) A	184) C	185) C	186) C	187) A	188) A	189) A	190) B
191) B	192) B	193) D	194) A	195) C	196) C	197) D	198) C	199) A	

NEET PREVIOUS YEARS QUESTIONS-ANSWERS

- 2 (c) 3 (a) 4 (a) 5 (d) 6 (a) 7 (a) 8 (a) 9 (a) 10 (d)
- 11 (b) 12 (b) 13 (c) 14 (c) 15 (c) 16 (b) 17 (a) 18 (a) 19 (a) 20 (c)
- 21 (b) 22 (b) 23 (a) 24 (a) 25 (a) 26 (a) 27 (c) 28 (a) 29 (b) 30 (2)
- **31** (4) **32** (3) **33** (1) **34** (3) **35** (2) **36** (3) **37** (4) **38** (3) **39** (3) **40** (1)
- **41** (2) **42** (4) **43** (1) **44** (1) **45** (1) **46** (2) **47** (4) **48** (2) **49** (1) **50** (1)
- **51** (1) **52** (2) **53** (3) **54** (3) **55** (1) **56** (1) **57** (4) **58** (2) **59** (1)

NEET PREVIOUS YEARS QUESTIONS-EXPLANATIONS

- 1. (b) Semi-conservative DNA replication was first shown in bacterium Escherichia coli by Matthew Meselson and Franklin Stahl.
- 2. (c)
- 3. (a) A polyribosome (or polysome) is a complex of an mRNA molecule and two or more ribosomes that act to translate mRNA instructions into polypeptides.
- **4. (a)** Coding strand and mRNA have the similar nucleotide sequence except, Thymine (T) is replaced by Uracil (U) in mRNA.
- **5.** (d) Unlike eukaryotes, in prokaryotes the genes are organised into operon. Operon is a co-ordinated group of genes which are all transcribed together & regulate a metabolic pathway as a unit.
- 6. (a) 7. (a)
- **8.** (a) Hershey and Chase proved that DNA is genetic material. They used bacteriophage for their experiment.
- **9. (a)** DNA fragments are negatively charged because of presence of phosphate group.

- **10.** (d) Ribosomal RNA (rRNA) is most abundant in animal cell. It constitutes about 80% of total RNA of the cell.
- **11. (b)** The association of H1 protein indicates the complete formation of nucleosome which requires DNA condensation. Therefore DNA is in condensed form.
- **12. (b)** If deletion happens at 901st position than the remaining 98 bases specifying for 33 codons of amino acids will be altered.
- 13. (c)
- **14.** (c) In eukaryotes spliceosomes are used in removal of introns during post-transcriptional processing of hnRNA. They are absent in prokaryotes.
- 15. (c)
- **16. (b)** Zinc-finger analysis is used for protein analysis. The zinc finger proteins are a super family of proteins involved in numerous activities of plant growth and development.
- 17. (a) The start codon is the first codon of a messenger RNA (mRNA) transcript translated by a ribosome. The start codon always codes for methionine in eukaryotes and a modified Met (fMet) in prokaryotes. The most common start codon is AUG.
- **18.** (a) A polysome or polyribosome is a complex of an mRNA molecule and two or more ribosomes, which is formed during the active translation process. They were initially named as ergosomes in 1963. However, further research by Jonathan Warner and Alex Rich characterised polysome.
- 19. (a)
- **20.** (c) Chargaff's rule is not applicable to RNA.
- 21. (b) 22. (b)
- 23. (a) Lac operon under control of repressor shows a negative regulation. Operon has inducible nature.
- 24. (a)
- **25.** (a) Synthesis of RNA exhibits several features that are synonymous with DNA replication. RNA synthesis requires accurate and efficient initiation, elongation proceeds in the 5'–3' direction (*i.e.* the polymerase moves along the template strand of DNA in the 3'–5' direction), and RNA synthesis requires distinct and accurate termination. Transcription exhibits several features that are distinct from replication.
- 26. (a) 27. (c) 28. (a)
- **29. (b)** Human genome sequencing is a process that determines the complete DNA sequence of an organism's genome at a single time. This requires sequencing of an organism's chromosomal DNA as well as DNA contained in the mitochondria and, for plants, in the chloroplast. Commonly used vectors for human genome sequencing are BAC (Bacterial artificial chromosomes) and YAC (Yeast artificial chromosomes).
- 45. Opening of DNA helix is facilitated by RNA polymerase. Opening of DNA is performed by DNA helicase
- 46. In the first step of RNA translation mRNA binds to ribosomes
- 47. 6.6×10^9 bp x 0.34 nm = 2.2 meters
- 48. DNA—DNA-----Replication

DNA---RNA---- Transcription

RNA—Protein ---- Translation

d—protein

- **49.** Split gene arrangement is characteristic of eukaryotes.
 - In capping 5-methyl guanosine triphosphate is added at 5' end of hnRNA.
 - At 3' end poly-A tail is added.
 - The non coding or template strand is copied to an mRNA. RNA polymerase associate with ρ factor (Rho factor) and it alters the specificity of the RNA polymerase to terminate the processes.
- **50.** RNA polymerase III transcribes tRNA, ScRNA, 5S rRNA and SnRNA.
 - RNA polymerase I transcribes 5.8S, 18S and 28S rRNA.
 - RNA polymerase II transcribes hnRNA which is precursor of mRNA
- The enzyme capable of Initiation , elongation and termination in the process of transcription in prokaryotes
 - DNA dependent RNA polymerase

- 52. Palindromic Nucleotide sequences are specific recognition sequence
- 53. Short interfering RNA or silencing RNA is not involved in protein synthesis.
- 54. The codon 'AUG' codes for methionine and 'UUU' codes for phenylalanine.
- * Histones are rich in basic amino acids residue lysine and arginine with charged side chain.
 - * There are five types of histone proteins i.e., H1, H2A, H2B, H3 and H4. Four of them occur in pairs to produce a unit of 8 molecules (histone octamer)
 - * The pH of histones is basic.
- 56. Repetitive DNA
- 57. z, y, a genes will be transcribed normally as i gene gets mutated and its product cannot bind the inducer molecule
- 58. Length of DNA is 2.2m i.e., $6.6 \times 10^9 bp$ Therefore 1.1m is $3.3 \times 10^9 bp$
- 59. 3 cycles of division occurs in 60 minutes forming a total of 80 *E.coli* cells. 20/80 cells contains medium DNA and 60/80 cells contain light DNA

